6719968

09945182

FIGURE 1

COMPARISON OF HUMAN V1-1 VS. HUMAN MP-52

V1-1 Ser Arg Cys Ser Arg Lys Pro Leu His Val Asp Phe Lys Glu Leu 1 AGC CGC TGC AGC CGC AAG CCG TTG CAC GTG GAC TTC AAG GAG CTC MP52 GCT CGC TGC AGT CGG AAG GCA CTG CAT GTC AAC TTC AAG GAC ATG 1 Ala Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp Asp Asp Trp Ile Ile Ala Pro Leu Asp Tyr Glu Ala Tyr GGC TGG GAC GAC TGG ATC ATC GCG CCG CTG GAC TAC GAG GCG TAC GGC TGG GAC GAC TGG ATC ATC GCA CCC CTT GAG TAC GAG GCT TTC 16 Gly Trp Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe 31 His Cys Glu Gly Leu Cys Asp Phe Pro Leu Arg Ser His Leu Glu CAC TGC GAG GGC CTT TGC GAC TTC CCT TTG CGT TCG CAC CTC GAG 31 His Cys Glu Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu 46 Pro Thr Asn His Ala Ile Ile Gln Thr Leu Leu Asn Ser Met Ala 121 CCC ACC AAC CAT GCC ATC ATT CAG ACG CTG CTC AAC TCC ATG GCA Pro Thr Asn His Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Asp Ala Ala Pro Ala Ser Cys Cys Val Pro Ala Arg Leu Ser CCA GAC GCG GCG CCC TCC TGC TGT GTG CCA GCG CGC CTC AGC 181 Pro Glu Ser Thr Pro Pro Thr Cys Cys Val Pro Thr Arg Leu Ser . 76 226 Pro Ile Ser Ile Leu Phe Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val Glu Ala Cys Gly Cys Arg AAG CAA TAC GAG GAC ATG GTG GTG GAG GCC TGC GGC TGC AGG 271 AAG CAG TAT GAG GAC ATG GTC GTG GAG TCG TGT GGC TGC AGG Lys Gln Tyr Glu Asp Met Val Val Glu Ser Cys Gly Cys Arg

Homology at the nucleotide level: 249/312 = 79.8%Homology at the amino acid level: 84/104 = 80.8%

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